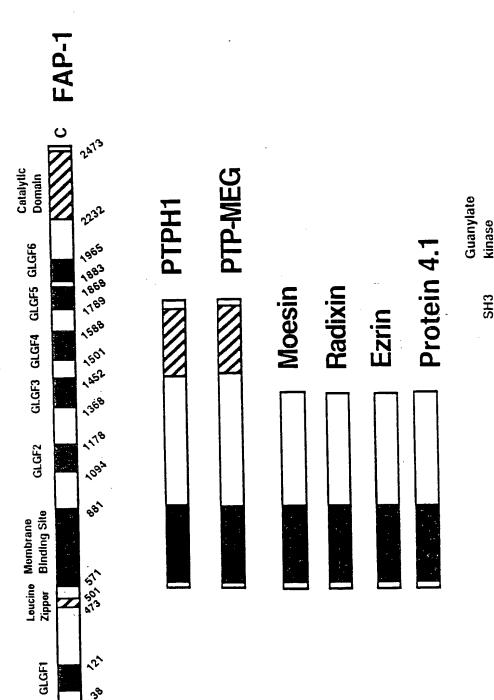
Z

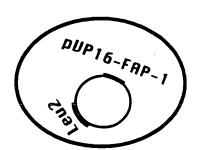


SH3 Kinase PSD-95

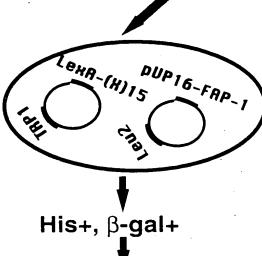
2/28 FIG. 2A

Construction of pBTM116 (LexA)-(X)15

Library DNAs of pBTM116 (LexA)-(X)15



Large scale transformation of yeast L40



**Curing of pVP16-FAP-1** 

Isolation of pBTM116 (LexA)-(X)15

Analysis of DNA sequences

ഗ Z FIG. 2C

FIG. 2D

NENEGOCIL

G

0 0

S

Mouse

ഗ o G Ш

SNSBBBB

S

Rat

ш

z

Z S

z

ш

D S

FIG. 2B Human

3/28

8-1

PIPIDISIEIDIGINIEIEIQIS

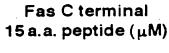
RSQLASV

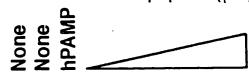
DSEMYNF

•									
12-0	2-0	13-0	20-0	6-2	9-2	18-1	22-1	71-1	14-5
>	>	>	>	>	>	>	>	>	_
	ш	а	O			Z	Z	A	1
		Н							
5	S	0	0	S	Р	>	>	≥	ш
<b>5</b>	G V S	F		SH	П	<u>۸</u>	Р	<b>%</b> >	Ш
A	5	A	O	Ш		S	Я		Z
$\vdash$	H	H	H	H	H	H		H	
A	A	5	A	S	Я		スス	¥	æ
<b>X</b>	N N	3	王	S		S	Y	r G K	ASR
CYA	Ш	W M G	EHA	N S S	75	GSD		-	A

ldogelwe czeci

FIG. 3A





200 -

97.4 -

69 -

FAP-1

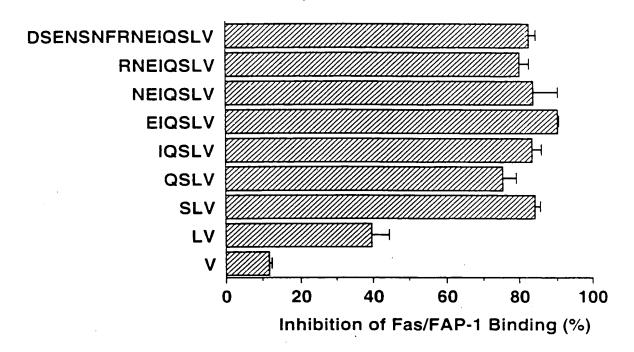
46 -

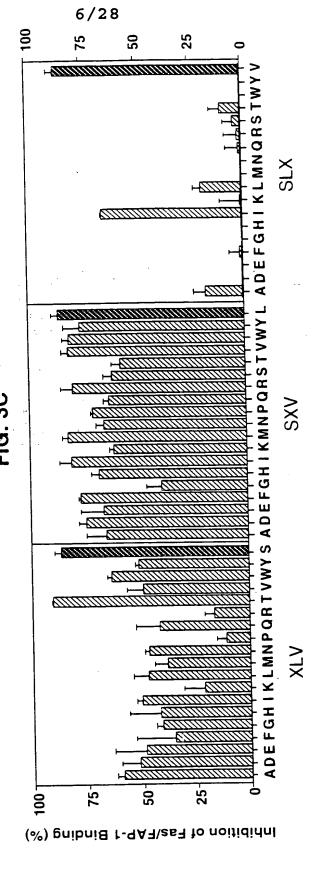
**30** –

21.5 **–** 14.3 **–** 

6 7 8 9 10

FIG. 3B





#### FIG. 4A

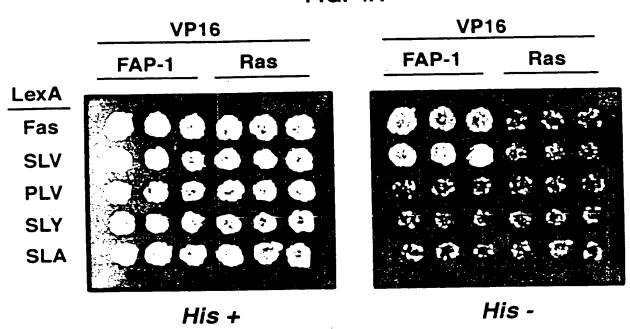
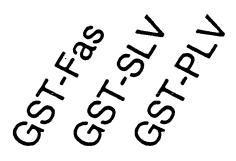


FIG. 4B



250 **-**148 **-**

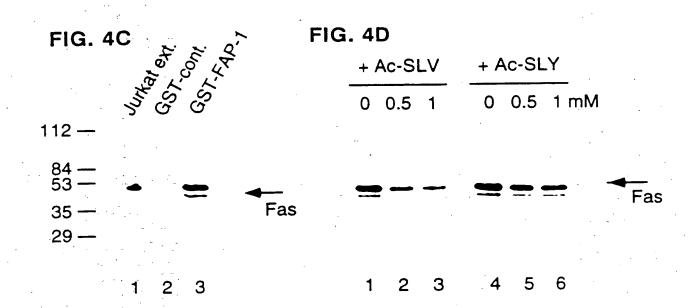


60 -

42 -

30





Phase contrast

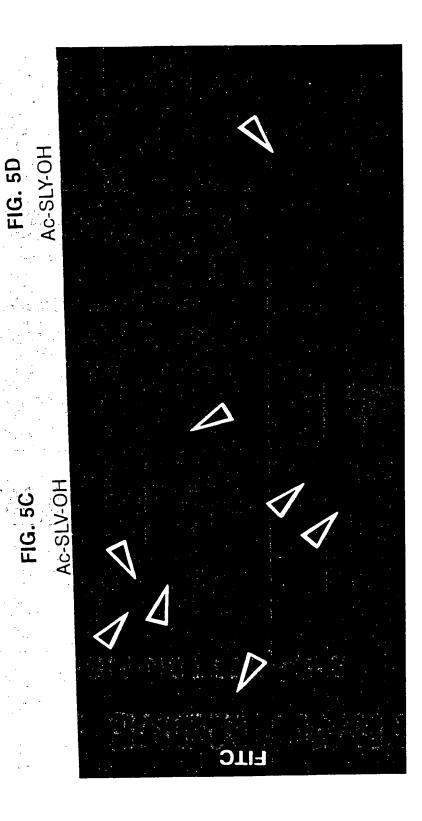
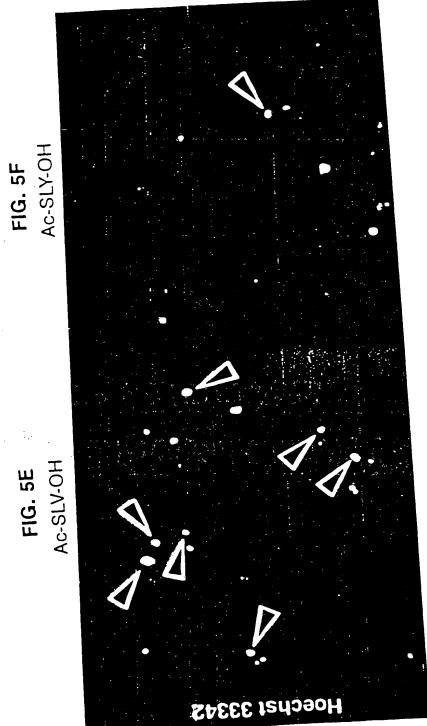
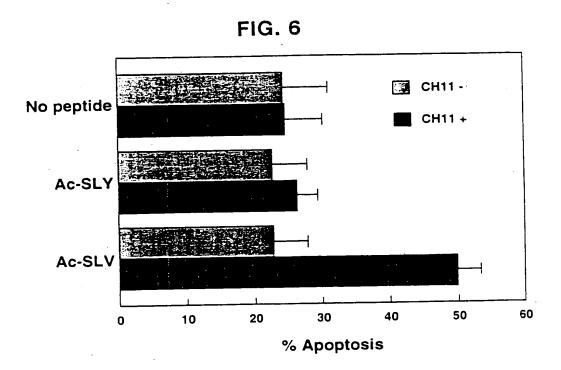


FIG. 5E





#### FIG. 7A

## NGF Receptor

nkggansrpv ngtpppegek adlveslcse lngsagdtwr egvaqpcgan ygyyddettg edterqlrec vvttvmgssq llaalrrigr acptglyths gecckacnlg pakreevekl eaddavcrca qdliastvag vdpclpctvc dgtysdeanh stdepeappe afkrwnsckg gdgglysslp atqdsatlda gldsmsapcv cpvrallasw lgvslggake epckpctecv tgtasggalk kqntvceecp ppegsdstap avvvglvayi sgslhdqqph lipvycsila gsglvfscqd ipgrwitrst vtfsdvvsat dgpr111111 qtvcepclds mgagatgram trwadaecee pvvtrgttdn **lhsdsgisvd** hlagelgyqp rceacrvcea statspv 61 181 241 301

#### FIG. 7B

### CD4 Receptor

ctasgkksig fhwknsngik kltgsge.lww yagsgnltla vskrekavwv figlgiffcv vsqlelqdsg iedsdtyice vedgkeevgl hltlpgalpg fsfplaftve slklenkeak kniqggktls vlggvaglll dsrrslwdgg nfpliiknlk vykkegegve klqmgkklpl psvqcrsprg gkkgdtvelt esnikulptw stpvqpmali qlqknltcev wgptspklml faktc**spi** ekktcdcphr qaerasssks witfdlknke vsvkrvtqdp kkvefkidiv vlafqkassi ltlesppgss 1 mnrgvpfrhl llvlqlallp aatggkkvvl rcrhrrrqae rmsqikrlls kgpsklndra thllqgqslt evnlvvmrat 11sdsgqv11 1npeagmwqc ilgnggsflt lvfgltansd twtctvlqnq leaktgklhg 181 301 241

#### FIG. 7C

	_	
Species	C-terminal sequences of NGFR (p75)	Binding activity of FAP-1
Human	SESTATSPV-COOH	+
Rat	SESTATSPV-COOH	+
Chicken	SESTATSPV-COOH	+

rttcsenela nlvaayekak nvegrkkss wekelagire gpsspgrits **issig**vsssv agcsvqpwes esihidplsy streageday kklakaqceq selrselsgs ksqndlltit slilgqfraa gtererdlle ssdrpvlgse dgscggafav fyndlkrans erlnsriehl 111alaeseg gttireedey elgrvitgle ypnlaeersr qtrlqsvqat kekkalelkl aspalelael aavkltmlel rphtnett rialleeens seirhqqsae elkaqlylle dkpgkecada ysaqcieaye divelnkrlq skirefevet dyiqq1\mdr elnkkidrlg dadacsdins etegvlgrdl drlrrrvrel klisktreess naakallmkl hetgvrmlkg slsstssgsk 1piakiaerv lyshgsalse natairlalq rahdcrktae ftkedegrlk elmamkeema vsaleritks hcdlaiktve atmuaireer pengetmyta ritelhsvia hsaalaslkg esquanter skeeelnrtk msmlvgkyee gdenitamlk sstasscdte csniqeifqt ldlenavlmg qerttlryee hieglttase veeqkegrmr ndssaelsel mdqdqtsvs1 elstssssnd kklkarvqel aeftnairre mnsgvamkyg lvhiehlkse leecksnaer sphagespyt ssnahtst shlmrehedv ghevnedsrs cslsvaevdr enesitamic thrpinpstg ashlahslqd dvkprgdsgr 601 421 541 661 241 301 361 481 181

#### FIG. 7E

mlagqppfdg pegdeegnme drlyfvmeyv gkvmladrkg ldseghikia vrehaffrri sdfegfsyvn sprfcdhcgs eklhvtvrda 1kpsdkdrr1 shctdfiwgf eegeyynvpi g flmvlgkgsf tglhscfqtv pownesttfk wwaygvllye yrdlkldnvm riylkaevad asgwykling eldrykltdfn flalidkopfl tlflinkrgii ayqpygkavd glmtkhpakr cgmdhtekrg 1 ktktiretin 1 kgpdtddprs vhevkdhkf1 ipdpkneskq fgvselmkmp futfscpgad sedrkapsan vectavekrv lskeavsick vfyaaeisig kgaenfdkff arkgalrgkn tpdyiapeii kgcvinvpsl gpagnkvisp kkdvvigddd trndfmgsls qvgkfkepqa mehnvsypks dgvttrtfcg cfvvhkrche kedtedmnvh 1sdpyvk1k1 appfkpkvcg tasqdvanrf gkagfacavc (11yglihagm ) teelyaikil nggdlmybiq dweklenrei pdfvhpilds dfgmckehmm ededelfqsi madvfpgnds sveiwdwdrt kn] ipmdpng lrqkfekakl 241 301 361 421 181 541 601

#### FIG. 7F

vntipalayk lekklqnatn yflmslaiad asimbleais Idryvaiqup vixegsella ddnivligsi seklfqrsih dafnwtvdse nrtnlscego faflpgesle sitrimavick ndcsmvalgk qhseeaskdn sdgvnekvgo fsrylqcqyk enkkplqlil wlyldvlfst e pvfglgddsk v glgtraklas f 1fvvmwcpff gnilvimavs vityfltiks lqkeatlevs tmgsisneqk ackvlgivff lssavnplvy tlfnktyrsa wplpsklcav eavviiltia t1svg1smp1 Inddtrlysn smltilygyr kaflkiiavw lestinslmq lgekmwsall lspsclellh mllgflvmpv vsffipleim repgaytgrr md:lceents ihnsrfnsrt 301

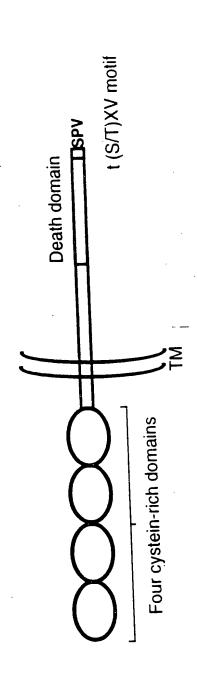
#### FIG. 7G

reskiyfrnp gdkteeqvex kalpnøgdet dscnqttlgm tafikituw asmivtyflt alltinfeam vamldgsrkd atkevktlrk dtllltene iganqynera fithitivic lvglfvmp1 gslaafftpl vdryiaikkp kerfgdfmlf detpcsspek flfllmwcpf fgryitenyr raskvlgivf tlfnktfrda stfvhvissn lekklqyatn asimhleais npnniccvlt vknkppgrlt witvetvfgr hgirnginpa kevatieneg wlfldvlfst vasgvnplvy gntlvilavs sigiatpy. lplvlcpa mrrtstigk maenskffkk ihalqkkayl 181 241 301 361 421

#### FIG. 7H

```
1 maaasydq11 kqvealkmen snlrqeledr snhltklete asnmkevlkq lqqsiedeam
    il assggidle rikelnidss nfpgvklrsk msirsygere gsvssrsged spvpmasipr
   121 rgfvngsres tgyleeleka rsliladidk eekekdwyya qiqmitkrid sipitenisi
   181 qtdmtrrqle yearqirvam eeqlqtcqdm ekraqrriar iqqiekdilr irqllqsqa:
   241 eaerssqnkh eigshdaerd negdgygein matagngggs tirmdhetas vissssthsa
   301 preltahlgt kvemvyslls mlgthdkddm artliamses odscismrqs golpliiqll
   361 hgndkdsvll gmsrgskear arasaalhni ihsqpddkrg rreirvihll eqiraycett
   421 wewqeahepg mdqddcpmpa pvehqicpav cvlmklsfde ehrhamnelg glqaiaellq
   481 vdcemygltn dhysitlrry agmaltnitf gdvarkatic smkgcmralv agiksesedi
541 qqviasvirn iswradvnsk ktirevgsvk almecalevk kestiksvis alwnisahct
    601 enkadicavd galaflygtl tyrsqtntla iiesgggilr nyssliatne dhrqilrenn
   661 clqtllqhlk shsltivsna cgtlwnlsar npkdqealwd mgavsmlknl ihskhkmiam
    721 gsaaairnim anrpakykda nimspgssip sihvrkqkai eaeldaqhis etfdnichis
  781 pkashrekqr hkqslygdyv fdtnrhddnr sdnfntgnmt vlspylnttv lpssassrqs
  841 ldasrsekdr slerergigl gnyhpatenp gtsskrolgi sttaaglakv meevsaihts
  901 qedrssgstt elhevtdern alrrssaaht hshtynftks enshrtesmi yakleykrss
961 ndslnsvsss dgygkrgenk psiesysedd eskfesyggy padlahkihs animidnidge
1021 ldtpinyslk ysdeglnsgr gspsenerwa rpkhiledel kesegresin esttypyyte
   1081 stddkhlkfq phfgqqecvs pyrsrgangs etnrvgsning inqnvsqslc qeddyeddkp
1141 tnyserysee eqheeeerpt nysikyneek rhvdqpidys lkyatdipss qkqsfsfsks
   1201 saggaskteh masssentst pasnakrong lhpssagsra gopokaatek vasingetig
1261 tycvedtpic farcaslasi saaedeigen ottgeadsan tigiaeikek igtraaedpv
   1321 sevpavsqmp rikasrlqqs slssesarhk avefssgaks paksgaqtpk sppehyvqet
   1381 plmfarctsv ssldsfesrs lassvqsepc sgmvsgilsp sdlpdspgqt mppsraktpp
   1441 pppqtaqtkr evpknkapta ekresgpkqa avnaavqrvq vlpdadtilh fatestpdqf
   1501 scssslsals ldepfickdv elrimppvce ndngmetese opkesnence keaektidse
1561 kdilddøddd dieileecii samptkesrk akkpaqtask lpppvarkps glpvykllps
1621 cmrlqpckhv sftpgddmpr vycvegtpin fstatsledl tiesppnela agegvrggaq
1681 sgefekrdti ptegrstdea oggktssvti pelddnkaee gdilaecins ampkgkshkp
1741 frykkimday agasasssap nkaqldakkk kotspykpip anteyrtryr knadskanla
   1801 aervisdikd skkonlknis kdindklinin edrvigsiai dspinytpie gtpycisind
1861 slssldiddd dvdlsrekae likakenkes eakvishtel tsnqqsankt qalakopini
    1321 gapkpilaka stfpasski pargaatdek lanfalentp vofshnssls sladidaenn
1381 nkenepiket eppdsageps kpassyspk sfhvedtpvc farnsslasi sidseddlla
2041 ecissampkk kkparlkadn ekhapringa ilgeditldl kdiarpaseh glapdsenfa
    2101 wkaigegans ivssihqaaa aacisrgass dsdsiislks gislgspfhl tpdqeekpft
    2161 snkaprilkp gekstletkk ieseskaika akkvykslit akvrsnseis gamkapiqan
    2221 mpsisrgrum ihipgvinss sstspvskkg pplktpasks psegqtatts prgakpsvks
2281 elepvarqts qiggsskaps rsqsrdstps rpaqqplsrp iqspgrnsis pgrnqisppn
    2341 klsqlprtss petastkasg sgkmaytapg romagqaltk qtglakmass ipraesaskg
2401 lnommignga nkkvelarma stkasgaesd raerpylvrq stflkeapap tlrrkleesa
    2461 efeslapesr pasptragag tpvlspslpd malathasvq aggwrklppm leptieymdg
    2521 rpakridiar shaespsilp inregtwhie hakhasalpr vetwritgsa asilsasses
     2581 sekaksedek hymsisgtko skenovsako twrkikenef spinstsotv ssgaingaes
     2641 ktliygmapa vsktedvævr ledopinnpr sgrsptgntp pvidsvseka npnikdskdn
     2701 qakqnvgngs vpmrtvglen rlnsfiqvda pdqkgteikp gqnnpvpvse imessivert
     2761 pissesskh espegivaar vipinynpep rkssadsisa rpsqipipvm nnikkrdekt
     2821 datessgtqs pkrhsqsylv tex
```

# (Low-affinity nerve growth factor receptor) p75NGFR



	C-terminal amino acid sequence
Fas	NEIQSLV
p75NGFR	STATSPV

PDZ domain t (S/T)-X-V |-COOH

interaction

In vitro interaction of 35S-labeled FAP-1 with various receptors FAP-1 binds to the cytoplasmic region of p75NGFR.

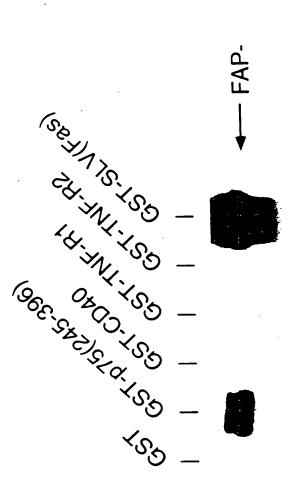
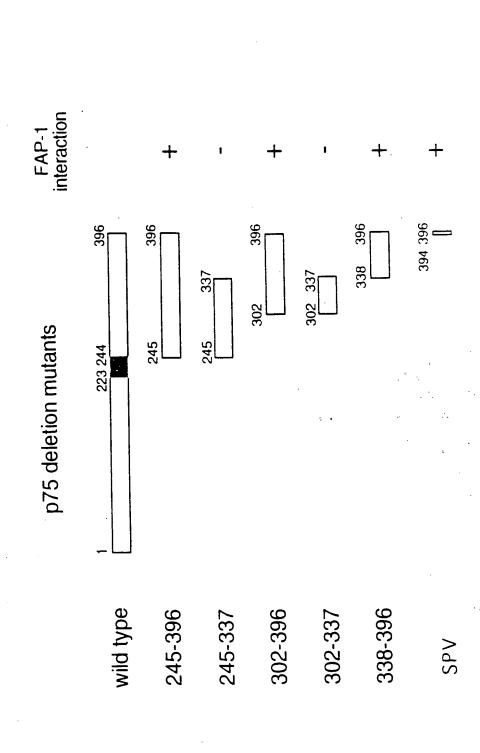


FIG. 11A

FAP-1 binds to C-terminal three amino acids SPV of p75NGFR.



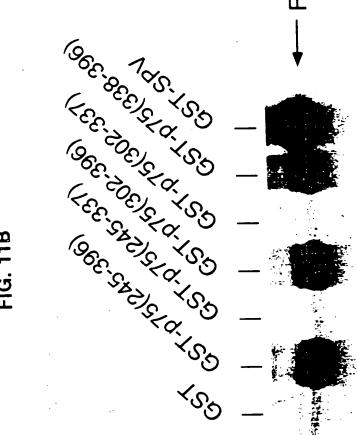
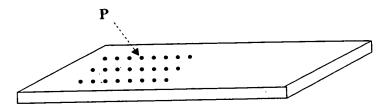


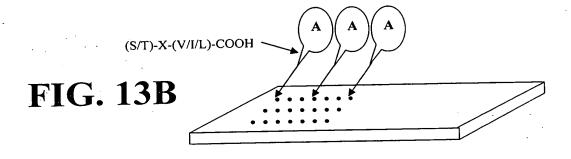
FIG. 12

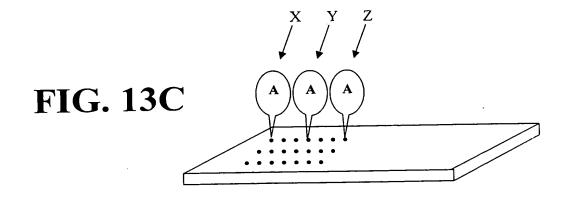
# FAP-1 binds to p75NGFR C-terminal cytoplasmic region in yeast.

	VP16-FAP-1	VP16-cRaf
LexA-p75NGFR(338-396)	+	•
LexA-p75NGFR(365-396)	+	ı
LexA-Fas	++	•
LexA-Ras <sup>V12</sup>	•	+
LexA-Lamin	•	







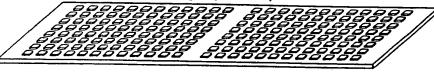


#### **FIG. 14A**

Plain-glass slide

**FIG. 14B** 

3D gel pad chip



**FIG. 14C** 

Microwell chip

